

Comparison of the immunogenicity of the human papillomavirus (HPV)-16/18 vaccine and the HPV-6/11/16/18 vaccine for oncogenic non-vaccine types HPV-31 and HPV-45 in healthy women aged 18–45 years

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Key words: *Cervarix*[®], cervical cancer, cross-protection, *Gardasil*[®], HPV-31, HPV-45, human papillomavirus, immunogenicity

Abbreviations: AAHS, amorphous aluminum hydroxyphosphate sulphate; ANOVA, analysis of variance; AS04, adjuvant system 04; ATP, according-to-protocol; CI, confidence interval; CIN, cervical intraepithelial neoplasia; cLIA, competitive luminex immunoassay; ED₅₀, effective dose producing 50% response; ELISA, enzyme-linked immunosorbent assay; GM, geometric mean; GMR, geometric mean (titer) ratio; GMT, geometric mean titer; GSK, GlaxoSmithKline; HPV, human papillomavirus; MPL, monophosphoryl lipid A; PBNA, pseudovirion-based neutralization assay; SD, standard deviation; TLR-4, toll-like receptor 4; TVC, total vaccinated cohort; VLP, virus-like particle

Protection against oncogenic non-vaccine types (cross-protection) offered by human papillomavirus (HPV) vaccines may provide a significant medical benefit. Available clinical efficacy data suggest the two licensed vaccines [HPV-16/18 vaccine, GlaxoSmithKline Biologicals (GSK), and HPV-6/11/16/18 vaccine, Merck and Co., Inc.,] differ in terms of protection against oncogenic non-vaccine HPV types -31/45. The immune responses induced by the two vaccines against these two non-vaccine HPV types (cross-reactivity) was compared in an observer-blind study up to Month 24 (18 mo post-vaccination), in women HPV DNA-negative and seronegative prior to vaccination for the HPV type analyzed [HPV-010 (NCT00423046)]. Geometric mean antibody titers (GMTs) measured by pseudovirion-based neutralization assay (PBNA) and enzyme-linked immunosorbent assay (ELISA) were similar between vaccines for HPV-31/45. Seropositivity rates for HPV-31 were also similar between vaccines; however, there was a trend for higher seropositivity with the HPV-16/18 vaccine (13.0–16.7%) vs. the HPV-6/11/16/18 vaccine (0.0–5.0%) for HPV-45 with PBNA, but not ELISA. HPV-31/45 cross-reactive memory B-cell responses were comparable between vaccines. Circulating antigen-specific CD4+ T-cell frequencies were higher for the HPV-16/18 vaccine than the HPV-6/11/16/18 vaccine [HPV-31 [geometric mean ratio (GMR) = 2.0; p = 0.0002] and HPV-45 [GMR = 2.6; p = 0.0092]], as were the proportion of T-cell responders (HPV-31, p = 0.0009; HPV-45, p = 0.0793). In conclusion, immune response to oncogenic non-vaccine HPV types -31/45 was generally similar for both vaccines with the exception of T-cell response which was higher with the HPV-16/18 vaccine. Considering the differences in cross-protective efficacy between the two vaccines, the results might provide insights into the underlying mechanism(s) of protection.

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Introduction

Worldwide, nearly 530,000 new cases of cervical cancer are diagnosed annually, and 274,000 women die of the disease each year.¹ Persistent infection with human papillomavirus (HPV) is the necessary cause for cervical cancer. Among the 15 oncogenic types identified, HPV-16 and HPV-18 are the most common and cumulatively account for approximately 70% of invasive cervical cancer cases.^{2,3} HPV-45 and -31 are the third and fourth most common oncogenic types globally, contributing to an additional 7% of cervical cancer cases.⁴ In addition to HPV-16 and -18, HPV-45 is associated with adenocarcinoma of the cervix, which is more difficult to detect.⁴

Two prophylactic HPV vaccines are currently licensed and available in more than 100 countries: *Cervarix*[®] (HPV-16/18 vaccine GlaxoSmithKline, Rixensart, Belgium) and *Gardasil*[®] (HPV-6/11/16/18 vaccine Merck, Whitehouse Station NJ USA). Both vaccines consist of virus-like particles (VLPs) of recombinant L1 capsid proteins from HPV-16 and -18. The HPV-6/11/16/18 vaccine also contains L1 VLPs from non-oncogenic types HPV-6 and -11, which are implicated in 75–90% of genital warts.^{5–7} The VLPs in the HPV-16/18 vaccine are formulated with the proprietary Adjuvant System 04 [AS04; 3-*O*-desacyl-4'-monophosphoryl lipid A (MPL) and aluminum hydroxide], which has been shown to induce higher cellular and humoral immune responses than VLPs formulated with aluminum salt alone.^{8,9} The VLPs in the HPV-6/11/16/18 vaccine are formulated with a proprietary amorphous aluminum hydroxyphosphate sulfate (AAHS) adjuvant.^{10,11} Both vaccines have demonstrated protection against infections and pre-cancerous lesions associated with oncogenic types HPV-16/18 in large, multicenter, randomized clinical trials.^{12–19}

In an early study, the HPV-16/18 vaccine demonstrated broad protection against histopathological outcomes beyond that anticipated for HPV-16/18, and also against incident infection with HPV-31/45.¹⁵ This led to further investigation into the protection offered by the HPV-16/18 vaccine against non-vaccine oncogenic HPV types (cross-protection). The cross-protective efficacy of the HPV-16/18 vaccine was further demonstrated in the large Phase III clinical efficacy study (HPV-008; PATRICIA, NCT00122681) using both virological and histopathological endpoints.¹⁷ In the according-to-protocol cohort for efficacy (ATP-E; all evaluable women who received three vaccine doses, had normal or low-grade cytology at baseline, and were evaluable for efficacy), vaccine efficacy was 77.5% [96.1% confidence interval (CI): 68.3–84.4] against 6-mo persistent HPV-31 infection and 92.0% (66.0–99.2) against HPV-31 associated cervical intraepithelial neoplasia (CIN) 2+ lesions. Vaccine efficacy was 76.1% (59.1–86.7) against 6-mo persistent HPV-45 infection and 100% (–67.8–100) against HPV-45 associated CIN2+ lesions.¹⁷ Cross-protection against HPV-31/45 was also shown in the total vaccinated naive cohort (TVC-naive) and the total vaccinated cohort (TVC).¹⁷ Cross-protection was also observed against persistent infection and CIN2+ associated with HPV-33.¹⁷

Cross-protection has also been evaluated in efficacy studies with the HPV-6/11/16/18 vaccine. It should be noted that these

results cannot be directly compared with those for the HPV-16/18 vaccine due to differences in study design, immunological assays, endpoints and study populations. In a cohort of women who were HPV DNA negative for each of the 4 vaccine types and 10 non-vaccine HPV types, protection against HPV-31 was 46.2% (95% CI: 15.3–66.4) for persistent infection and 56.9% (28.6–74.8) for CIN1–3 or adenocarcinoma in situ. For the other types evaluated, including HPV-45, efficacy was not statistically significant.²⁰

We previously presented results from study HPV-010 (NCT00423046), a comparative immunogenicity and safety study of the HPV-16/18 vaccine and the HPV-6/11/16/18 vaccine up to Month 24.^{21,22} At Month 7, the HPV-16/18 vaccine induced significantly higher anti-HPV-16/18 type-specific neutralizing antibody levels and significantly higher frequencies of HPV-16/18 type-specific circulating memory B cells than the HPV-6/11/16/18 vaccine. We now present data from a sub-analysis comparing the immune response induced by the HPV-16/18 vaccine and the HPV-6/11/16/18 vaccine against the oncogenic non-vaccine HPV types -31 and -45 (cross-reactivity). These immunological investigations were performed to provide insights into the potential underlying mechanism(s) of clinical cross-protection against HPV-31 and -45.

Results

Study population. A total of 1,106 women (553 women in each vaccine group) stratified into three age groups (18–26 y, 27–35 y and 36–45 y) were vaccinated as described in Einstein et al. 2009.²² The ATP cohort for immunogenicity included all subjects who were HPV DNA-negative and seronegative prior to vaccination for the HPV type under analysis, received three vaccine doses and for whom immunogenicity endpoints were available (671 women at Month 24: 341 women in the HPV-16/18 vaccine group and 330 women in the HPV-6/11/16/18 vaccine group).

Serum antibody responses. Seropositivity rates (Fig. 1) and geometric mean titers (GMTs) of neutralizing antibodies against HPV-31 and HPV-45 (Fig. 2) were measured by PBNA in the ATP cohort for immunogenicity, which was stratified into three age groups: 18–26 y, 27–35 y and 36–45 y. Across all age cohorts, one month after completion of the three-dose vaccination course (Month 7), 58.4–80.4% of women in the HPV-16/18 vaccine group were seropositive (defined as an antibody titer ≥ 40 ED₅₀) for HPV-31 compared with 31.4–46.2% of women in the HPV-6/11/16/18 vaccine group; 18.8–27.3% of women in the HPV-16/18 vaccine group and 3.1–9.5% of women in the HPV-6/11/16/18 vaccine group were seropositive for HPV-45 (Fig. 1). At Month 24, 10.5–38.9% of women in the HPV-16/18 vaccine group compared with 26.1–33.3% of women in the HPV-6/11/16/18 vaccine group remained seropositive for HPV-31; in the HPV-16/18 vaccine and HPV-6/11/16/18 vaccine groups, 13.0–16.7% and 0.0–5.0% of women, respectively, remained seropositive for HPV-45 (Fig. 1). As anticipated and across all age groups, GMTs peaked at Month 7 for HPV-31 (HPV-16/18 vaccine: 102–206 ED₅₀; HPV-6/11/16/18 vaccine, 41–50 ED₅₀) and for HPV-45 (HPV-16/18 vaccine, 29–33

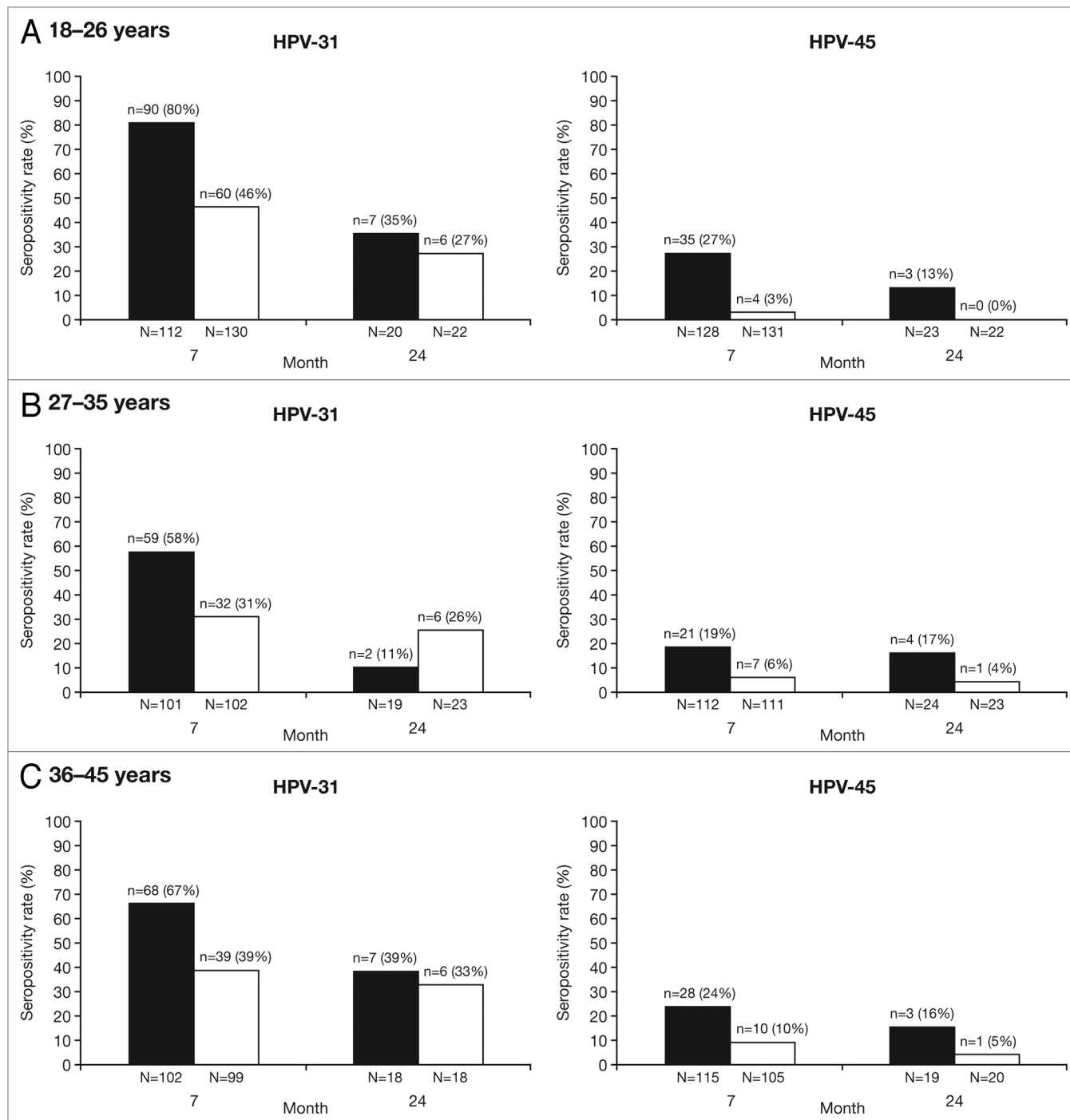


Figure 1. Seropositivity rates for anti-HPV-31 and anti-HPV-45 serum neutralizing antibodies measured by pseudovirion-based neutralization assay at Months 7 and 24 (ATP cohort for immunogenicity, seronegative and DNA-negative at baseline for HPV type analyzed). N, number of evaluable subjects; n, number of seropositive subjects, per vaccine, per timepoint. Percentages indicate seropositivity rates at each timepoint [seropositivity defined as neutralizing antibody titer ≥ 40 ED₅₀ (the PBNA limit of quantification)]. Black bars, Human Papillomavirus Bivalent (Types 16 and 18) Vaccine (Recombinant, adjuvanted, adsorbed) (*Cervarix*); white bars, Human Papillomavirus Quadrivalent (Types 6, 11, 16 and 18) Vaccine, Recombinant (*Gardasil*). The ATP cohort for immunogenicity included all evaluable subjects who received three vaccine doses (i.e., those meeting all eligibility criteria and complying with the procedures defined in the protocol) for whom data concerning immunogenicity endpoint measures were available.

ED₅₀; HPV-6/11/16/18 vaccine, 21–24 ED₅₀). With both vaccines and across all age groups, HPV-31 PBNA GMTs decreased from Month 7 to Month 12 and plateaued through Month 24 to levels close to or below the assay cut-off (40 ED₅₀); at Month 24, neutralizing antibody levels were 27–49 ED₅₀ for the HPV-16/18 vaccine and 35–38 ED₅₀ for the HPV-6/11/16/18 vaccine. HPV-45 neutralizing antibody GMTs decreased from Month 7 to Month 12 and then plateaued through to Month 24 (24–27 ED₅₀ for the HPV-16/18 vaccine and 20–21 ED₅₀ for the HPV-6/11/16/18 vaccine); at all timepoints, HPV-45 neutralizing antibody levels were below the PBNA limit of detection (Fig. 2).

Seropositivity rates and vaccine-induced serum antibody titers against HPV-31/45, assessed by ELISA in the ATP cohort for immunogenicity are shown in **Figures 3 and 4**, respectively. At Month 7, seroconversion rates were similar for both vaccines against HPV-31/45 across all age groups (97.4–100% for the HPV-16/18 vaccine vs. 100% for the HPV-6/11/16/18 vaccine). At Month 24, 66.7–85.6% of women in the HPV-16/18 vaccine group and 64.0–68.9% in the HPV-6/11/16/18 vaccine group remained seropositive for HPV-31; 68.9–89.0% of women in the HPV-16/18 vaccine group compared with 73.7–78.2% of women in the HPV-6/11/16/18 vaccine group remained seropositive for HPV-45 (Fig. 3). GMTs in women aged 18–26 y were generally higher for the HPV-16/18 vaccine than the HPV-6/11/16/18 vaccine for both HPV-31 (215 and 109 ELISA units/mL, respectively) and HPV-45 (193 and 114 ELISA units/mL, respectively), and were similar with either vaccine in women aged 27–35 and 36–45 y (Fig. 4).

Memory B-cell responses. HPV cross-reactive memory B-cell responses were evaluated in a subset of women in the ATP cohort for immunogenicity who had no detectable B-cell response prior to vaccination. The geometric means (GMs) of circulating antigen-specific memory B-cell responses were measured only in ‘responders’ (defined as subjects with detectable HPV cross-reactive memory B-cells, i.e., ≥ 1 specific memory B-cell/million memory B-cells; see methods) because data for all subjects did not follow a normal distribution. The highest percentage of memory B-cell responders was observed at Month 7 for HPV-31 (HPV-16/18 vaccine, 76.0%; HPV-6/11/16/18 vaccine, 67.3%; $p = 0.3838$) and HPV-45 (HPV-16/18 vaccine, 63.5%; HPV-6/11/16/18 vaccine, 64.9%; $p = 1.0000$). At Month 24, there were approximately equal proportions of responders between vaccine groups for both HPV-31 (HPV-16/18 vaccine, 63.6%; HPV-6/11/16/18 vaccine, 62.9%; $p = 1.0000$) and HPV-45 (HPV-16/18 vaccine, 51.4%; HPV-6/11/16/18 vaccine, 50%; $p = 1.0000$) (Fig. 5).

Cross-reactive memory B-cell GMs at Month 24 (calculated only in responders) were comparable for both groups for HPV-31 and -45 (Fig. 6). GM ratios (GMRs) of the response to the HPV-16/18 vaccine vs. the HPV-6/11/16/18 vaccine for HPV-31 and -45 were 0.69 (95% CI: 0.41, 1.14; $p = 0.1438$) and 0.95 (0.55, 1.62; $p = 0.8426$), respectively.

CD4+ T-cell responses. Cross-reactive CD4+ T-cell responses were evaluated in a subset of women in the ATP cohort for immunogenicity who were T-cell-negative prior to vaccination

(<500 HPV cross-reactive CD4+ T-cells/million CD4+ T-cells at baseline). The proportion of T-cell responders [defined as subjects with ≥ 500 HPV cross-reactive CD4+ T-cells identified in vitro as expressing two or more of four immune markers (CD40L, IL-2, TNF α , IFN γ) per million cells] was analyzed between vaccine groups]. At Month 7, the proportion of T-cell responders were similar between vaccines for HPV-31 (HPV-16/18 vaccine, 85.4%; HPV-6/11/16/18 vaccine, 68.4%; $p = 0.1070$) and significantly higher for HPV-45 with the HPV-16/18 vaccine vs. the HPV-6/11/16/18 vaccine (76.6% vs. 51.2%, $p = 0.0223$). At Month 24, the proportion of T-cell responders was significantly higher in the HPV-16/18 vaccine group than the HPV-6/11/16/18 vaccine group for HPV-31 (86.7% vs. 43.3%, $p = 0.0009$), and higher in the HPV-16/18 vaccine group than the HPV-6/11/16/18 vaccine group for HPV-45 (62.5% vs. 37.5%, $p = 0.0793$) (Fig. 7). The GM of the frequency of circulating antigen-specific CD4+ T-cells in all subjects at Month 24 was significantly higher in the HPV-16/18 vaccine group than the HPV-6/11/16/18 vaccine group for both HPV-31 [GMR = 2.0 (GM: 813, 409); $p = 0.0002$] and HPV-45 [GMR = 2.6 (GM: 668, 257); $p = 0.0092$] (Fig. 8).

Discussion

The cross-protective efficacy of the HPV-16/18 vaccine against HPV-31/33/45 was demonstrated in an earlier large clinical study (HPV-008)^{17,23,24} in a cohort of women who were HPV DNA-negative for corresponding HPV type at baseline, regardless of serostatus.¹⁷ In the total vaccinated cohort for efficacy, vaccine efficacy against 6 mo persistent infection attributed to HPV-31, -33 and -45 increased over time from an interim analysis, performed at a mean follow-up of 14.8 mo [standard deviation (SD): 4.9 mo] after third vaccine dose (36.1%, 36.5% and 59.9%, respectively),²⁴ through to analysis at 34.9 mo (SD: 6.4 mo) (66.9%, 42.2% and 71.6%, respectively).¹⁷ Recent results from the end-of-study analysis of this trial confirmed the cross-protective efficacy of the HPV-16/18 vaccine against these three HPV types through to Month 48.²³ Together, HPV types -16, -18, -31, -33 and -45 account for approximately 82% of cervical cancers.²⁵

Cross-protection results have also been published for the HPV-6/11/16/18 vaccine,²⁰ although it should be noted that these should not be directly compared with the HPV-16/18 results, as the study designs of the HPV-16/18 vaccine and HPV-6/11/16/18 vaccine trials differ in HPV DNA- and immunological assays, endpoints and study populations. In a cohort of women who were seronegative and DNA-negative for HPV types in the HPV-6/11/16/18 vaccine, and DNA-negative for ten non-vaccine HPV types (including HPV-31/45), the HPV-6/11/16/18 vaccine demonstrated cross-protective efficacy against CIN1–3 or adenocarcinoma in situ associated with HPV-31; no cross-protection was shown against CIN1–3 or adenocarcinoma in situ associated with HPV-45.²⁰

The current study was designed to directly compare the immune response to non-vaccine oncogenic HPV types elicited by the HPV-16/18 vaccine and the HPV-6/11/16/18

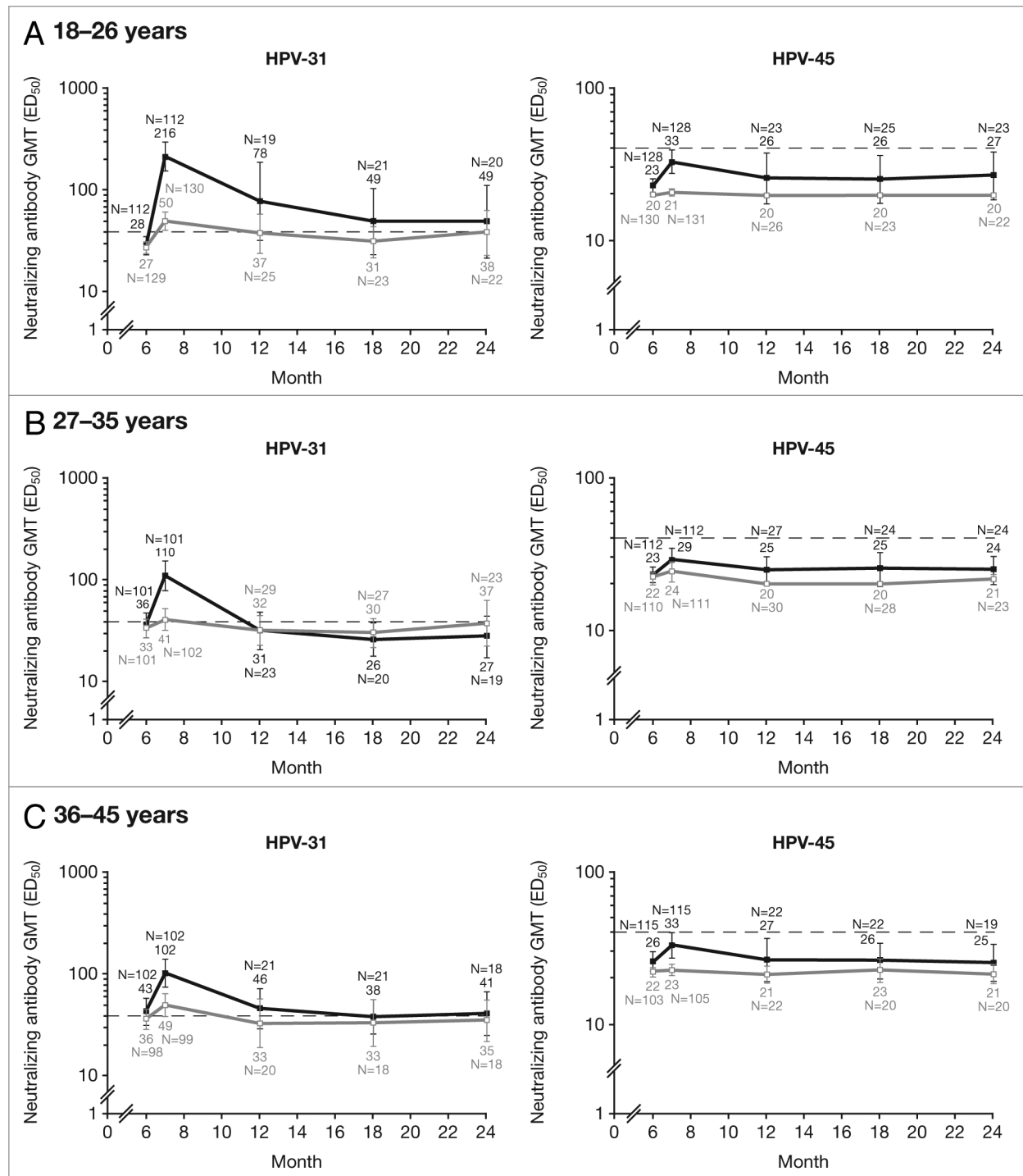


Figure 2. Geometric mean titers (GMTs) for anti-HPV-31 and anti-HPV-45 serum neutralizing antibodies measured by pseudovirion-based neutralization assay at Months 7, 12, 18 and 24 (ATP cohort for immunogenicity, seronegative and DNA-negative at baseline for HPV type analyzed). ED₅₀, effective dose producing 50% response; GMT, geometric mean titer. Black square line, Human Papillomavirus Bivalent (Types 16 and 18) Vaccine (Recombinant, adjuvanted, adsorbed) (*Cervarix*); white square line, Human Papillomavirus Quadrivalent (Types 6, 11, 16 and 18) Vaccine, Recombinant (*Gardasil*). Error bars denote 95% confidence intervals (CIs) of GMTs; dashed line, PBNA limit of detection (40 ED₅₀). N, number of evaluable subjects. The ATP cohort for immunogenicity included all evaluable subjects who received three vaccine doses (i.e., those meeting all eligibility criteria and complying with the procedures defined in the protocol) for whom data concerning immunogenicity endpoint measures were available.

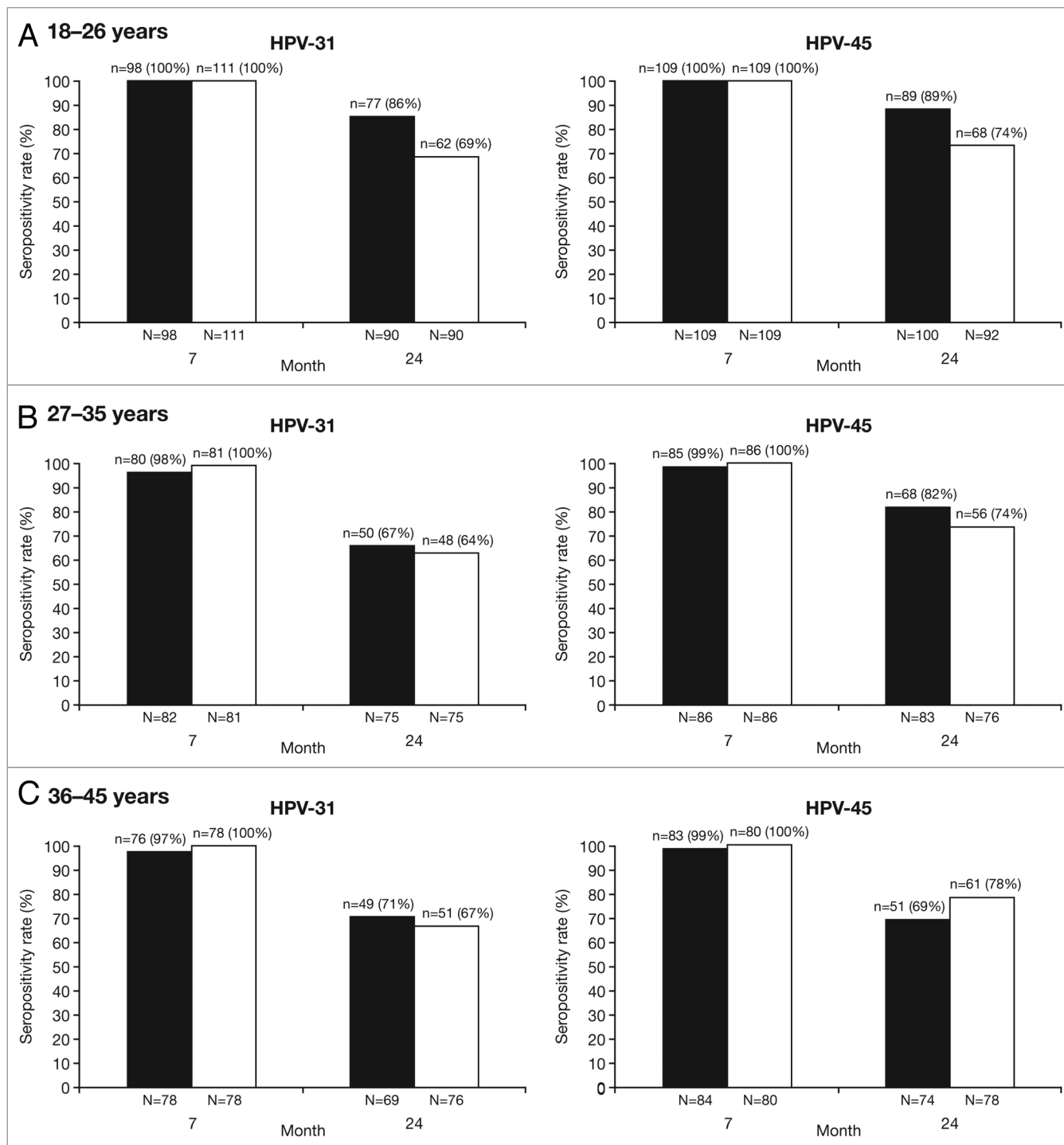


Figure 3. Seropositivity rates for anti-HPV-31 and anti-HPV-45 IgG antibodies measured by enzyme-linked immunosorbent assay at Months 7 and 24 (ATP cohort for immunogenicity, seronegative and DNA-negative at baseline for HPV type analyzed). N, number of evaluable subjects; n, number of seropositive subjects, per vaccine, per timepoint. Percentages indicate seropositivity rates at each timepoint (seropositivity defined as IgG titer ≥ 59 ELISA units/mL [the ELISA limit of quantification]). Black bars, Human Papillomavirus Bivalent (Types 16 and 18) Vaccine (Recombinant, adjuvanted, adsorbed) (*Cervarix*); white bars, Human Papillomavirus Quadrivalent (Types 6, 11, 16 and 18) Vaccine, Recombinant (*Gardasil*). The ATP cohort for immunogenicity included all evaluable subjects who received three vaccine doses (i.e., those meeting all eligibility criteria and complying with the procedures defined in the protocol) for whom data concerning immunogenicity endpoint measures were available.

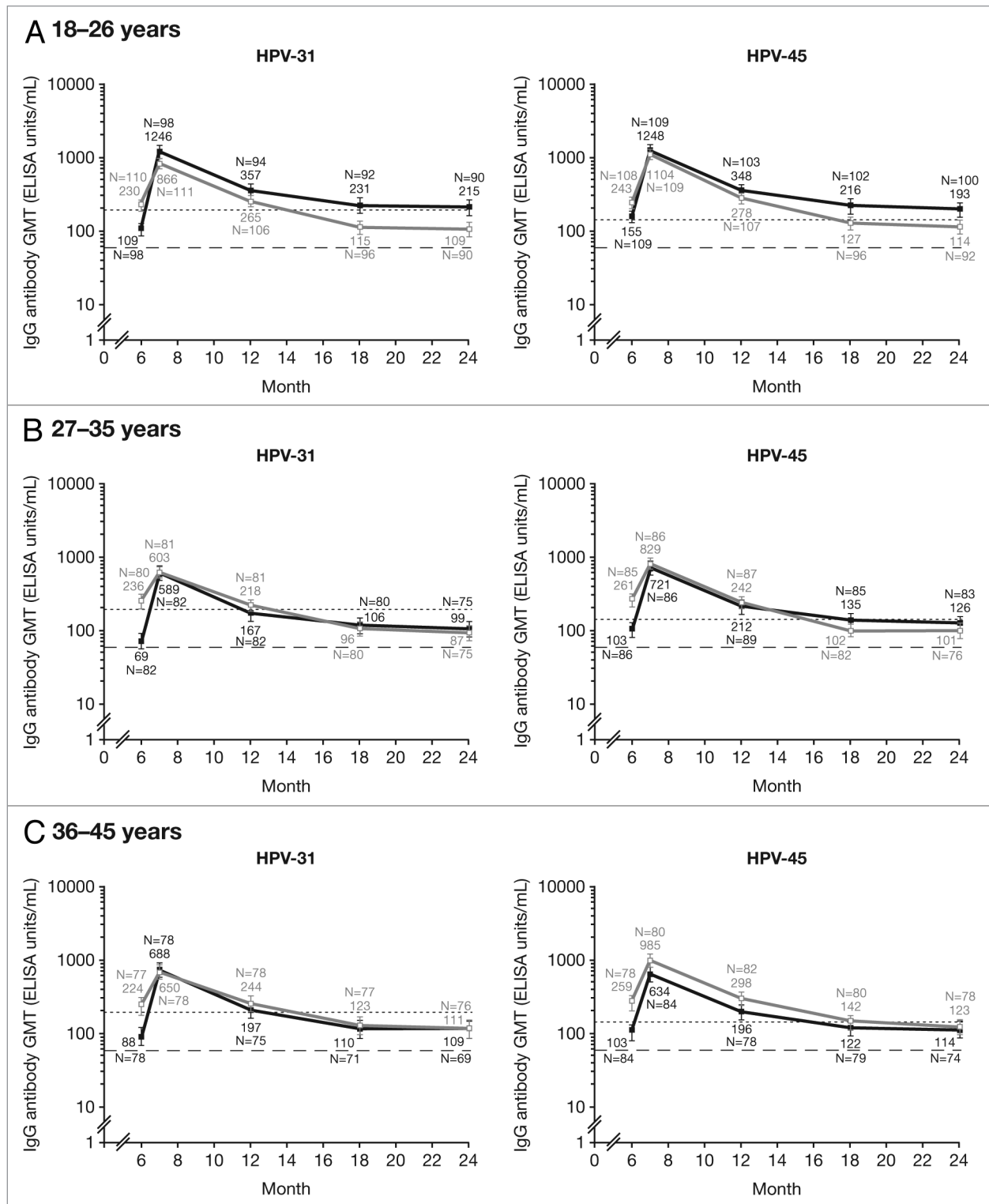


Figure 4. Geometric mean titers (GMTs) for anti-HPV-31 and anti-HPV-45 IgG antibodies measured by enzyme-linked immunosorbent assay at Months 7, 12, 18 and 24 (ATP cohort for immunogenicity, seronegative and DNA-negative at baseline for HPV type analyzed). GMT, geometric mean titer; IgG, immunoglobulin G. Black square line, Human Papillomavirus Bivalent (Types 16 and 18) Vaccine (Recombinant, adjuvanted, adsorbed) (Cervarix®); white square line, Human Papillomavirus Quadrivalent (Types 6, 11, 16 and 18) Vaccine, Recombinant (Gardasil®). Errors bars denote 95% confidence intervals (CIs) of GMTs. Dashed line, ELISA limit of detection (59 ELISA units/mL). Dotted line, GMTs for natural infection antibody levels (183.5 ELISA units/mL for HPV-31 and 139.0 ELISA units/mL for HPV-45). N, number of evaluable subjects. The ATP cohort for immunogenicity included all evaluable subjects who received three vaccine doses (i.e., those meeting all eligibility criteria and complying with the procedures defined in the protocol) for whom data concerning immunogenicity endpoint measures were available.

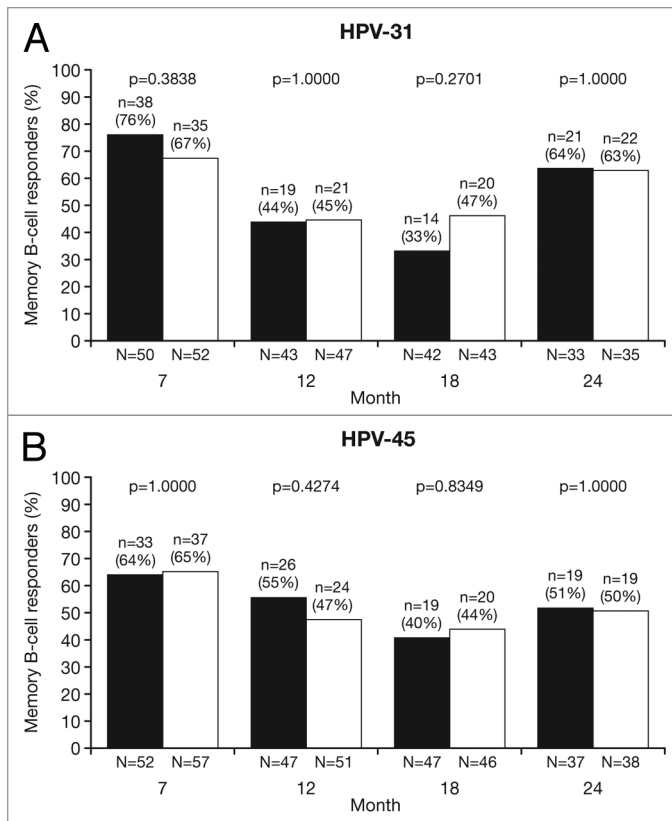


Figure 5. Proportion of responders for (A) HPV-31- and (B) HPV-45-specific memory B-cell responses at Months 7, 12, 18 and 24 (ATP cohort for immunogenicity; seronegative, DNA-negative and with no detectable HPV cross-reactive B-cells prior to vaccination). N, number of subjects with available results; n, number of responders, per vaccine, per timepoint; %, percentage of responders, per vaccine, per timepoint. Black bars, Human Papillomavirus Bivalent (Types 16 and 18) Vaccine (Recombinant, adjuvanted, adsorbed) (*Cervarix*[®]); white bars, Human Papillomavirus Quadrivalent (Types 6, 11, 16 and 18) Vaccine, Recombinant (*Gardasil*[®]). Responders defined as subjects with detectable HPV cross-reactive memory B cells (≥ 1 cell/million cells). p-values were calculated using Fisher's exact test to compare proportion of responders.

vaccine. Our sub-analysis of the HPV-010 study, along with the available data generated in clinical efficacy studies, was intended to provide insight into the potential mechanism(s) for cross-protection.

Our observations show that both the HPV-16/18 vaccine and the HPV-6/11/16/18 vaccine induced cross-reactive serologic responses against HPV-31 and HPV-45. Humoral immune responses for non-vaccine types HPV-31/45 (PBNA and ELISA) were much lower than those for vaccine types HPV-16/18.^{21,22} At Month 24, there were no significant differences between the HPV-16/18 vaccine and the HPV-6/11/16/18 vaccine in terms of HPV-31/45 circulating antibodies as measured by PBNA. It should be noted that circulating neutralizing antibodies were at levels close to or below the limit of detection of each assay. Similar levels of circulating antibodies (measured by ELISA) were observed between both vaccines in the 27–35 y and 36–45 y groups; for both vaccines, the highest levels of HPV-31 and HPV-45 circulating antibodies were reported in the 18–26 y group.

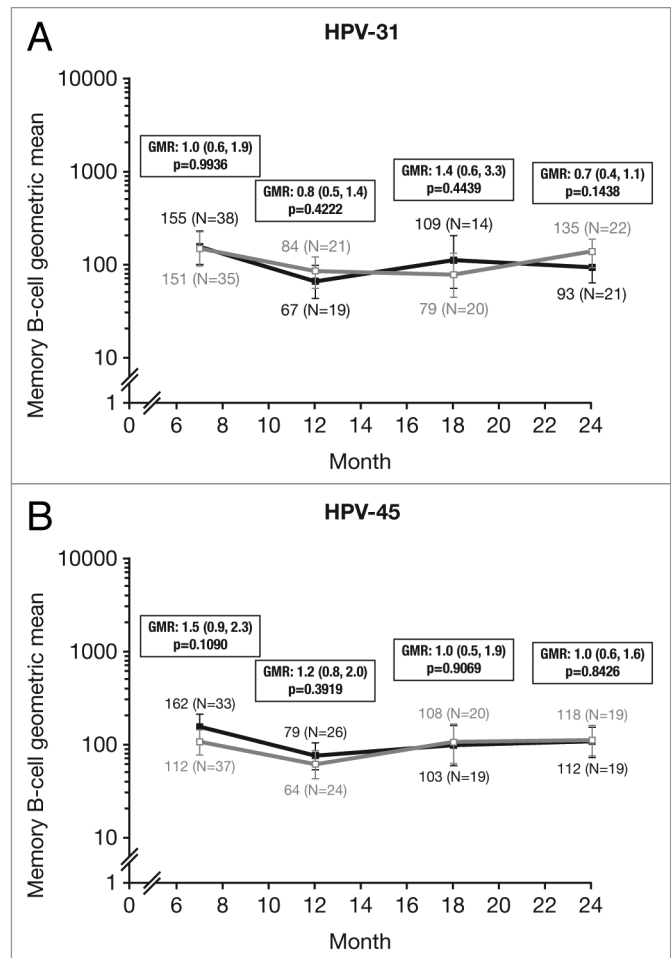


Figure 6. Geometric means (GM) and GM ratios (GMR) in responders only for (A) HPV-31- and (B) HPV-45-specific memory B-cells at Months 7, 12, 18 and 24 (ATP cohort for immunogenicity; seronegative, DNA-negative and with no detectable HPV cross-reactive B cells prior to vaccination). GMR, geometric mean ratio; N, number of responders [i.e., subjects with detectable HPV cross-reactive memory B cells (≥ 1 cell/million cells)]. Black square line, Human Papillomavirus Bivalent (Types 16 and 18) Vaccine (Recombinant, adjuvanted, adsorbed) (*Cervarix*[®]); white square line, Human Papillomavirus Quadrivalent (Types 6, 11, 16 and 18) Vaccine, Recombinant (*Gardasil*[®]). Error bars denote 95% confidence intervals of geometric means. Statistical comparison (GMR ANOVA p-value) was performed on B-cell responders because data for all subjects in the subset did not follow a normal distribution.

Previous results from the HPV-010 study demonstrated higher vaccine specific immune responses with the HPV-16/18 vaccine compared with the HPV-6/11/16/18 vaccine, as supported by higher levels of specific neutralizing antibodies, seropositivity rates and a higher frequency of HPV cross-reactive memory B cells and T cells.^{21,22} However, in our sub-analysis, the levels of neutralizing antibodies to HPV-31/45 were relatively low for both vaccines. There was also no significant difference in circulating HPV cross-reactive memory B-cell response for HPV-31 and HPV-45 with both vaccines, likely due to the fact that most memory B cells would have migrated to central lymphoid organs, such as the spleen, to be subsequently activated by the interaction of circulating antigen presenting cells such as CD4⁺ T cells.²⁶

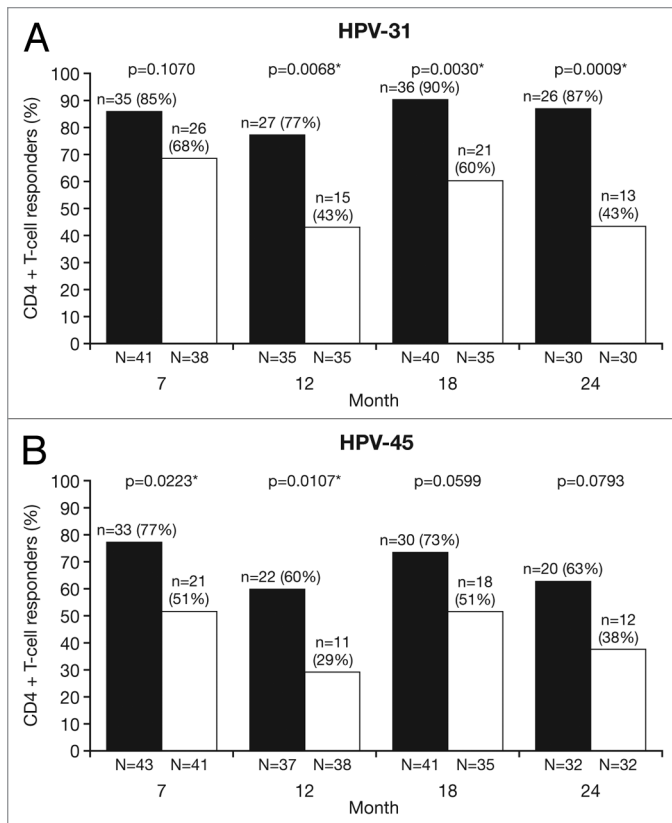


Figure 7. Proportion of responders for (A) HPV-31- and (B) HPV-45-specific CD4+ T-cell response at Months 7, 12, 18 and 24 (ATP cohort for immunogenicity; seronegative, DNA-negative and with a HPV-specific CD4+ T-cell response below 500 cells per million cells prior to vaccination). N, number of subjects with available results; n, number of responders, per vaccine, per timepoint; %, percentage of responders, per vaccine, per timepoint. * $p < 0.05$. Black bars, Human Papillomavirus Bivalent (Types 16 and 18) Vaccine (Recombinant, adjuvanted, adsorbed) (Cervarix®); white bars, Human Papillomavirus Quadrivalent (Types 6, 11, 16 and 18) Vaccine, Recombinant (Gardasil®). Responders defined as subjects with ≥ 500 HPV cross-reactive memory CD4+ T-cells expressing at least two of four immune markers (CD40L, IL-2, TNF α , IFN γ) per million cells. p-values were calculated using Fisher's exact test to compare proportion of responders.

An earlier study from women immunized with the HPV-6/11/16/18 vaccine, investigating vaccine-induced antibody binding and neutralization with HPV-18/45 using PBNA, also demonstrated relatively low levels of vaccine-induced antibodies to cross-neutralize HPV-45 pseudovirions.²⁷ Findings from a recent study by Kemp et al. suggest that low levels of cross-neutralizing antibodies may contribute to the mechanism of cross-protection observed with the HPV-16/18 vaccine.²⁸ It may simply follow that the response to HPV-16 and -18 is higher due to the immunodominance of the type-specific epitopes, so only a relatively small proportion are shared with phylogenetic family members like HPV-31 and -45, respectively.

A separate study showed that a fourth dose of the HPV-16/18 vaccine induced an anamnestic response (evidenced by a rapid and strong increase of antibody titers) not only to the vaccine types HPV-16/18, but also to non-vaccine types HPV-31/45.²⁹

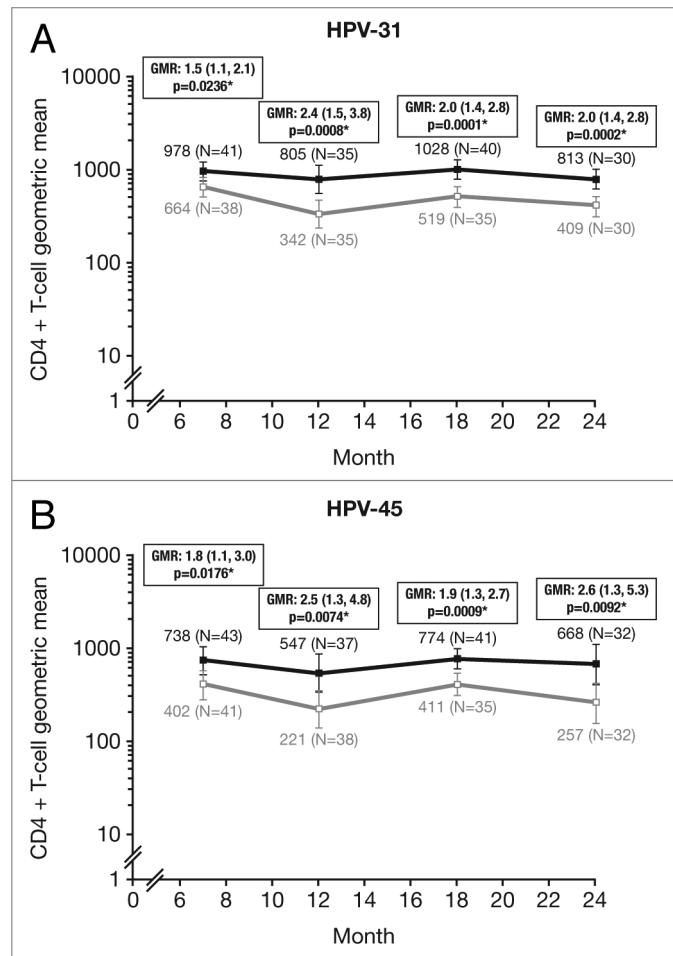


Figure 8. Geometric means (GM) and GM ratios (GMR) for (A) HPV-31- and (B) HPV-45-specific CD4+ T-cell response at Months 7, 12, 18 and 24 in all subjects in the subset (ATP cohort for immunogenicity; seronegative, DNA-negative and with a HPV cross-reactive CD4+ T-cell response below 500 cells per million cells prior to vaccination). * $p < 0.05$. GMR, geometric mean ratio; N, number of subjects with available results. Black square line, Human Papillomavirus Bivalent (Types 16 and 18) Vaccine (Recombinant, adjuvanted, adsorbed) (Cervarix®); white square line, Human Papillomavirus Quadrivalent (Types 6, 11, 16 and 18) Vaccine, Recombinant (Gardasil®). Error bars denote 95% confidence intervals of geometric means. Statistical comparison (GMR ANOVA p-value) was performed on all subjects.

The demonstration of immunologic memory for the HPV-31/45 response in women confirms that the HPV-16/18 vaccine does induce cross-reactive immune responses even though circulating neutralizing cross-reactive antibodies are at levels close to or below the limit of detection of the assays.³⁰

The main immunological parameter that differentiates the immune response to HPV-31/45 is the CD4+ T-cell response. In our sub-analysis, CD4+ T-cell responses remained high from Month 7 to Month 24 and the HPV-16/18 vaccine gave higher CD4+ T-cell responses than the HPV-6/11/16/18 vaccine at all timepoints. Importantly, in contrast to the relatively low levels of cross-reactive antibodies, the frequency and quality of the cross-reactive CD4+ T-cell responses to HPV-31 and -45 were similar to the specific responses to HPV-16 and -18.²² However, we note

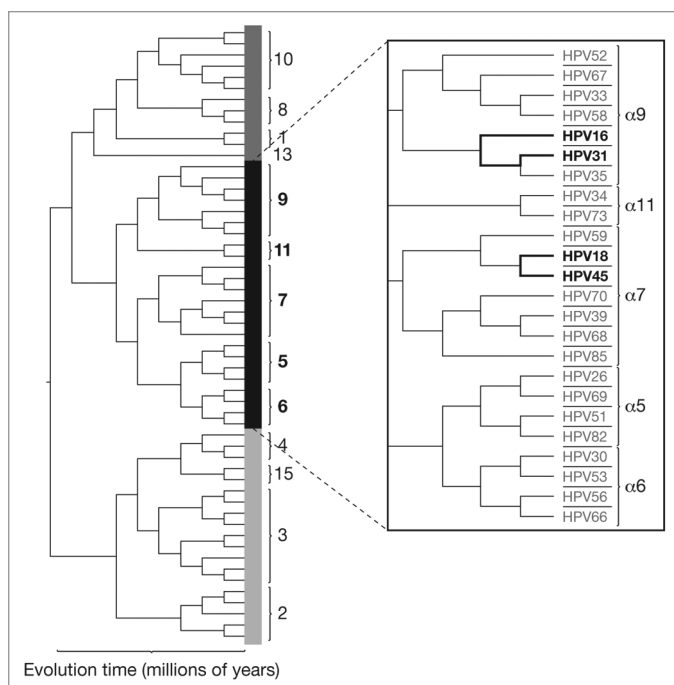


Figure 9. Phylogenetic tree of anogenital human papillomavirus types (adapted from Schiffman and Wentzensen 2010 and Schiffman et al. 2005).^{54,55} This phylogenetic tree is based on the alignment of concatenated early and late open reading frames. The carcinogenicity of HPV types reflects viral evolution. The clade presented in detail in the figure above (α5, 6, 7, 9, 11) reflects the HPV types associated with cervical cancers and their precursors.⁵⁴

that, based on published data,^{17,20} there appear to be differences in the HPV types for which the vaccines confer cross-protective efficacy. This suggests that higher levels of CD4⁺ T-cell response may be necessary for cross-protection against certain HPV types, such as HPV-45, but that the requirements for cross-protection against other types such as HPV-31 might be different.

Of note, the cross-reactive T-cell response to HPV-31 and -45 was measured using pools of synthetic peptides spanning the truncated VLP L1 sequences of HPV-31 and HPV-45. We cannot exclude the possibility that the HPV-6/11/16/18 vaccine induced a cross-reactive T-cell response against the portion absent (approximately 30 amino acids) from the truncated HPV-31/45 L1 VLPs. However, such bias is unlikely given that the truncated portion is small and also the similarity of the HPV-31/45 T-cell responses to the specific responses to HPV-16/18 (obtained using a pool of peptides spanning the entire VLP L1 sequences of HPV-16/18).

Other assays used in the current study are not anticipated to have introduced a bias toward either vaccine. As discussed previously in references 21 and 22, no bias was observed for HPV-16/18 data obtained using the ELISA or memory B-cell data obtained using the ELISPOT assay. For the PBNA used in the present study, the amino acid sequence of the L1 protein and the cell-line used for the production of the pseudovirions, which contain L1 and L2, are different from those used in either vaccine. Importantly, PBNA and ELISA have also been shown to correlate

with Merck's competitive Luminex immunoassay (cLIA).^{31,32} Moreover, none of the assays, nor the HPV-16/18 vaccine or the HPV-6/11/16/18 vaccine contain HPV-31/45 VLPs.

Adjuvant systems have been shown to enhance specific and cross-clade neutralizing antibody immunological responses in addition to T-cell responses.³³⁻³⁵ In the case of the HPV-16/18 vaccine, the AS04 adjuvant system,³⁶ which contains MPL (50 μg) absorbed on aluminum salt (Al³⁺, 500 μg), may similarly enhance the immune responses. The AS04 component of the HPV-16/18 vaccine was shown to induce a higher frequency of type-specific memory B cells and antibody responses against HPV-16/18 compared with the same vaccine formulated with aluminum salt alone.⁸ The MPL component of the AS04 adjuvant binds and activates the Toll-like receptor-4 (TLR-4), which is present on key antigen-presenting cells that play an important role in the induction of innate and adaptive immune responses.^{37,38} Furthermore, the combination of an aluminum salt with MPL is thought to prolong cytokine responses at the injection site.³⁶ Taken together, these factors may plausibly account for the higher CD4⁺ T-cell response against HPV-31/45 observed with the HPV-16/18 vaccine compared with the HPV-6/11/16/18 vaccine in this study.

Although the mechanism(s) of cross-protection has not yet been fully elucidated, it is likely to be associated with the phylogenetic relationship of HPV types. HPV-16, -18, -31 and -45 belong to the genus α-papillomavirus, which is further classified by species and then type; the A7 species includes HPV types -18 and -45 and the A9 species includes HPV types -16 and -31 (Fig. 9). HPV types belonging to the same species are phylogenetically related;^{39,40} based on predicted L1 amino acid sequences, HPV-31 shares 83% homology with the L1 protein of HPV-16 and HPV-45 shares 88% homology with the L1 protein of HPV-18.²⁰ The observed difference in the cross-protective efficacy for HPV-45 between the vaccines is probably not related to the primary sequence of the HPV-18 L1 VLPs, since both share a similar percentage of sequence homology with HPV-45. However, the truncation of ~30 amino acids at the C-terminus of the HPV-16/18 vaccine's HPV-18 L1 VLPs⁴¹ might impact on the accessibility of epitopes that are shared between HPV-18 and HPV-45.

Confirmation would require further biophysical structural characterization of HPV-18 L1 VLPs, the demonstration of differences in epitope recognition by antibodies induced by the two vaccines, or evaluation of the relative avidity of antibodies for the different epitopes. In addition, conformational differences in L1 VLP epitope exposure due to differences in vaccine production process and/or the adjuvant formulation (as discussed above) may also contribute to differences in the quality of cross-reactive neutralizing antibody responses, and may contribute to the differences in cross-protection observed between the vaccines.

Despite high levels of homology between HPV types of the same species, even if the conformational structures of L1 VLPs from different HPV types are very similar, the surface loops that contain neutralizing domains display significant amino acid heterogeneity. The unique features of these surface loops are the distinct surface immunodominant conformational epitopes

that provide type-specific protection. There are high levels of conserved homology between HPV types that may represent subdominant cross-reactive epitopes.⁴² As cross-neutralization induced by L1 VLPs represents less than 1% of the type specific neutralizing activity induced by the immunodominant conformational epitopes, it is uncertain whether this is sufficient to offer cross-protection *in vivo*.⁴³ The capacity of L1-VLP-induced antibodies to mediate type-specific and cross-protection against cervicovaginal challenge was recently demonstrated in a murine challenge model, following vaccination with HPV-16 VLPs (on alum) or following passive transfer of immune serum.⁴⁴ A high level of specific protection was observed against HPV-16 and partial cross-protection was observed against HPV-31 challenge.⁴⁴ Furthermore, in rabbits that received the HPV-16/18 vaccine, high levels of cross-protection against HPV-31/45 were observed after HPV quasivirion [virions with human HPV L1/L2 capsids and the cotton-tail rabbit papillomavirus (CRPV) genome, produced in 293TT cells]⁴⁵ challenge in the presence of no or low levels of neutralizing antibodies.⁴⁶ Importantly these controlled experimental studies clearly demonstrate the capacity of L1 VLPs to induce cross-protective immunity and indicate that the cross-protection observed in clinical studies resulted from vaccine-induced immune responses.

Investigations into the mechanism of L1 VLP-induced prevention of HPV infection have led to the proposal of two distinct mechanisms of protection by L1 specific polyclonal antibodies: (1) high levels of antibodies result in an immunoglobulin-coated HPV capsid, which prevents interaction of the capsid with the cell surface, (2) a lower antibody level allows the capsid to associate with the cell surface but prevents the conformational changes required for virus entry.^{44,47} It has been suggested that cross-protection and type-specific long-term protection may be attributed to the second mechanism with a relatively low antibody to virus ratio and that PBNA may not be a suitable assay to assess cross neutralizing antibodies.⁴⁴ The National Cancer Institute (United States) is currently investigating the possibility of developing more accurate methods to assess all sets of neutralizing antibodies.⁴⁷

An alternative role of antibodies in mediating cross-protection follows from the observation that HPV suppresses Langerhans cell activation, resulting in local immune suppression. Interestingly, this suppression can be reversed when Langerhans cells encounter HPV virions or L1/L2 VLPs in the presence of antibodies.^{48,49} Thus, HPV infection in the presence of vaccine-induced antibodies could activate Langerhans cells, via the Fc receptor or other mechanisms, and lead to local inflammation that does not occur in the absence of antibodies. Importantly, the antibodies would not need to have the functional capacity to neutralize the virus.

Furthermore, a role for CD4⁺ T-cells in cross-protection in the presence of cross-reactive antibodies is proposed. It has been shown that CD4⁺ T-cells can activate NK cells in the presence of pro-inflammatory signals,⁵⁰ suggesting that the combination of activated Langerhans cells, due to the presence of antibodies,⁴⁶ and elevated frequencies of cross-reactive cytokine producing CD4⁺ T-cells could activate IFN γ producing NK cell

responses.⁴⁸ Thus both CD4⁺ T- and NK cells could contribute to cytokine-mediated reduction of viral replication and/or elimination of viral infected epithelial cells.

Assuming that CD4⁺ T-cells and NK cells have a role to play, one would expect to see better cross-protection for progressive disease efficacy endpoints such as CIN2+/3+, due to elimination of infected cells, than for incident/persistent infection efficacy parameters. Although there is some evidence of this pattern being observed from AS04-adjuvanted HPV-16/18 vaccine efficacy studies, it is not unequivocal and further investigation will provide more complete data to verify this hypothesis.¹⁷

Notably, CD4⁺ T-cell responses have been shown to be important in limiting the progression of cervical cancerous lesions⁵¹ and in the regression of genital warts.⁵² In a study examining the efficacy of a novel therapeutic vaccine containing HPV-16 E6 and/or E7 synthetic peptides, 60% (95% CI: 36–81) of patients with HPV-16-positive vulvar intraepithelial neoplasia had a clinical response three months after last vaccination; patients with a complete response had a significantly higher CD4⁺ T-cell response than patients without a complete response.⁵¹ Therefore, a CD4⁺ T-cell-mediated mechanism that eliminates HPV-infected cells is plausible.

However, our study assessed only immunological endpoints; whether or not the higher levels of cross-reactive CD4⁺ T cells observed with the HPV-16/18 vaccine compared with the HPV-6/11/16/18 vaccine correlate with enhanced protection against the progression of cervical lesions has not been assessed.

In conclusion, vaccination with the HPV-16/18 vaccine or the HPV-6/11/16/18 vaccine induces humoral responses to non-vaccine HPV types -31/45, albeit at generally low levels; in the case of the HPV-16/18 vaccine, this response has demonstrated immunological memory in other investigations. The HPV-16/18 vaccine induced relatively higher HPV-31/45-specific CD4⁺ T-cell responses compared with the HPV-6/11/16/18 vaccine, which may play a role in cross-protection; however, further studies are necessary to fully understand and elucidate the roles (and possible interactions) of both humoral and cell-mediated immunity in the HPV-16/18 vaccine and the HPV-6/11/16/18 vaccine for protection against cervical lesions caused by oncogenic non-vaccine types. A limitation of our analysis is that it only evaluated the immune response to two oncogenic non-vaccine HPV types. The assessment of a greater number of non-vaccine types associated with cross-protection or a lack of cross-protection would be valuable to identify other immunological markers associated with cross-protection. This may increase our understanding of the underlying mechanism(s) of cross-protection, and with further follow-up, help to determine the duration of cross-protection and perhaps subsequently facilitate the determination of valid and universally accepted correlates of protection.

Materials and Methods

Study design. This was a sub-analysis performed at Month 24 of a Phase III trial conducted in 40 centers in the US (HPV-010; NCT00423046). The overall study design was described previously in references 21 and 22. Healthy women were stratified by

age (18–25, 27–35 and 36–45 y) and randomized in a 1:1 ratio to receive the HPV-16/18 vaccine or the HPV-6/11/16/18 vaccine according to their recommended three-dose schedules (Months 0, 1, 6 or Months 0, 2, 6, respectively). To ensure the study was conducted in an observer-blind manner, a dose of placebo (aluminum hydroxide) was given at either Month 1 or 2 as appropriate; the study is ongoing and will be maintained blind through 48 mo of follow-up. The study was conducted in accordance with the Declaration of Helsinki and Good Clinical Practice guidelines.

Blood sampling and assays. Analyses were performed on the ATP cohort for immunogenicity that included all women who met eligibility criteria, complied with the trial protocol, received all three doses of study vaccine and were DNA-negative and seronegative for HPV type considered at baseline, and for whom immunogenicity endpoint measures were available.

Blood sampling was performed as described previously in references 21 and 22. At Months 6, 7, 12, 18 and 24, additional samples were collected from a pre-specified subset of women in all age groups in both vaccine groups for additional immunological assessment, which included HPV cross-reactive memory B-cell and CD4⁺ T-cell responses. HPV-31/45 type-specific pseudovirion-based neutralization assays (PBNA), enzyme-linked immunosorbent assays (ELISA) and B-cell ELISPOT assays were performed in assays similar to those previously described for HPV-16/18.^{21,22} In brief, PBNA and ELISA were performed as described in Dessy et al.⁵³ with minor amendments as follows: (1) For PBNA, HPV-31 or -45 pseudovirions were used instead of HPV-16 or -18 pseudovirions; (2) for ELISA, HPV-31 and -45 VLP were used as coating antigen; instead of using eight two-fold serial dilution starting at 1/100, sera were diluted 1/100 and 1/1,000; the cut-off for ELISA was established as the concentration, in EU/mL, equivalent to the upper limit of the one-sided 95% confidence interval of a population of 18–25-y-old women (who had no more than one lifetime sexual partner, whose cervix was free of cytological abnormalities and who tested DNA negative for high-risk HPV by PCR and seronegative for HPV-16 and -18 by ELISA). For each antigen, seropositivity was defined as a neutralizing antibody titer ≥ 40 ED₅₀ (the assay threshold) for PBNA and an antibody titer ≥ 59 ELISA units/mL for ELISA. Responders for memory B cells were defined as subjects with detectable HPV cross-reactive memory B-cells (≥ 1 cell per million cells).

CD4⁺ T-cell responses to HPV-31/45 were evaluated in an assay similar to that previously described for HPV-16/18.²² Cross-reactive responses were evaluated in vitro by using a pool of 20-mer HPV peptides to stimulate peripheral blood antigen-specific CD4⁺ T-cells to produce cytokines. The 20-mer HPV synthetic peptides overlapped by ten amino acids and spanned the entire truncated sequences of the HPV-31/45 L1 VLPs. Intracellular cytokine staining, followed by flow cytometry, was used to quantify the number of cells producing immune markers (CD40L, IL-2, TNF α and IFN γ) in response to in vitro stimulation with four different peptide pools.²² Responders were defined as women who had ≥ 500 specific CD4⁺ T cells per million identified as producing two or more of the immune markers after in vitro stimulation.

Statistical analysis. The primary exploratory objective of the HPV-010 study was to compare HPV-16/18 antibody titers induced by the HPV-16/18 vaccine and the HPV-6/11/16/18 vaccine in women 18–26 y of age as measured by PBNA at Month 7.²¹ Results from Month 24 have also been published in reference 22. Secondary objectives included evaluating serum immune responses to HPV-31/45 in the ATP cohort for immunogenicity of women aged 18–26, 27–35 and 38–45 y of age at Months 6, 7, 12, 18 and 24 by PBNA and ELISA. Secondary objectives also included evaluating the frequency of HPV-31/45-specific memory B-cells and cytokine-positive CD4⁺ T-lymphocytes in a pre-specified 30% of women in each vaccine group. Statistical analyses (within-group and between-group comparisons) of immunogenicity were performed at Months 7, 12, 18 and 24, as described previously in reference 21.

Memory B-cell and CD4⁺ T-cell analysis. Memory B-cell and CD4⁺ T-cell responses for each stimulant at each timepoint were summarized for each group, HPV type, and age strata by the geometric mean (GM). In an exploratory analysis, the proportion of responders in each group was tested using Fisher's exact test.

The GMs of circulating antigen-specific memory B-cell responses were calculated only for subjects with no response at baseline and a response (≥ 1 cell/million cells) at the specific timepoint under analysis. The GMs were calculated only in responders as data for all subjects did not follow a normal distribution (samples with no detectable response were assigned a value of 1 for the purpose of GM calculation).

The GMs of circulating antigen-specific CD4⁺ T cells were assessed in all subjects (i.e., responders and non-responders). The GMR is the ratio between the GMs of vaccine groups. Both the ANOVA (analysis of variance) model and Kruskal-Wallis test were used to calculate p-values associated to GMRs; the ANOVA p-values are presented here.

Notes

Cervarix is a registered trade mark of the GlaxoSmithKline group of companies. *Gardasil* is a registered trade mark of Merck and Co., Inc.

Conflict of Interest Statement

All authors have completed the Unified Competing Interest form at www.icmje.org/coi_disclosure.pdf and declare: The submitted work got financial support from GlaxoSmithKline Biologicals.

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